AMENDMENTS TO THE CLAIMS

This Listing of Claims will replace all prior versions, including listings, of claims in the

application.

Listing of Claims

Claim 1 (original): An isolated peptide selected from the group consisting of:

(a) a peptide set forth in Tables 1-14; and

(b) a derivative of the peptide in (a).

Claim 2 (original): The isolated peptide of claim 1, wherein Xaa1 is Glu or γ-carbox y-Glu,

Xaa2 is Gln or pyro-Glu, Xaa3 is Pro or hydroxy-Pro, Xaa4 is Trp or bromo-Trp, and Xaa5 is Tyr,

¹²⁵I-Tyr, mono-iodo-Tyr, di-iodo-Tyr, O-sulpho-Tyr or O-phospho-Tyr.

Claim 3 (original): The derivative of the peptide of claim 1, in which the Arg residues may

be substituted by Lys, ornithine, homoargine, nor-Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-

trimethyl-Lys or any synthetic basic amino acid; the Lys residues may be substituted by Arg,

ornithine, homoargine, nor-Lys, or any synthetic basic amino acid; the Tyr residues may be

substituted with meta-Tyr, ortho-Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-

phospho-Tyr, nitro-Tyr or any synthetic hydroxy containing amino acid; the Ser residues may be

substituted with Thr or any synthetic hydroxylated amino acid; the Thr residues may be substituted

with Ser or any synthetic hydroxylated amino acid; the Phe residues may be substituted with any

synthetic aromatic amino acid; the Trp residues may be substituted with Trp (D), neo-Trp, halo-Trp

(D or L) or any aromatic synthetic amino acid; the Asn, Ser, Thr or Hyp residues may be

glycosylated; the Tyr residues may also be substituted with the 3-hydroxyl or 2-hydroxyl isomers

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(meta-Tyr or ortho-Tyr, respectively) and corresponding O-sulpho- and O-phospho-derivatives; the

acidic amino acid residues may be substituted with any synthetic acidic amino acid, e.g., tetrazolyl

derivatives of Gly and Ala; the aliphatic amino acids may be substituted by synthetic derivatives

bearing non-natural aliphatic branched or linear side chains C_nH_{2n+2} up to and including n=8; the Leu

residues may be substituted with Leu (D); the Glu residues may be substituted with Gla; the Gla

residues may be substituted with Glu; the N-terminal Gln residues may be substituted with pyroGlu;

the Met residues may be substituted by NIe; the Cys residues may be in D or L configuration and may

optionally be substituted with homocysteine (D or L); and pairs of Cys residues may be replaced

pairwise with isoteric lactam or ester-thioether replacements, such as Ser/(Glu or Asp), Lys/(Glu or

Asp), Cys/(Glu or Asp) or Cys/Ala combinations.

Claim 4 (withdrawn): An isolated nucleic acid encoding an conotoxin propeptide having an

amino acid sequence set forth in Table 1.

Claim 5 (withrdrawn): The isolated nucleic acid of claim 4, wherein the nucleic acid

comprises a nucleotide sequence set forth in Table 1.

Claim 6 (original): An isolated conotoxin propertide having an amino acid sequence set

forth in Table 1.

Claim 7 (withdrawn): A method of alleviating pain in an individual which comprises

administering to said individual that is either exhibiting pain or is about to be subjected to a

pain-causing event a pain-alleviating amount of an active agent comprising a pain-relieving

conotoxin peptide of claim 1 or a pharmaceutically acceptable salt thereof.

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Claim 8 (withdrawn): A method for treating or preventing disorders associated with a

disorder selected from the group consisting of voltage-gated ion channel disorders, ligand-gated ion

channel disorders and receptor disorders in an individual which comprises administering to an

individual in need thereof a therapeutically effective amount of a conotoxin peptide of claim 1 or a

pharmaceutically acceptable salt thereof.

Claim 9 (withdrawn): A method of identifying compounds that mimic the therapeutic

activity of a conotoxin, comprising the steps of: (a) conducting a biological assay on a test compound

to determine the therapeutic activity; and (b) comparing the results obtained from the biological

assay of the test compound to the results obtained from the biological assay of a conotoxin.

Claim 10 (original): A substantially pure conotoxin peptide derivative comprising a

permutant of the peptide of claim 1.

Claim 11 (original): A substantially pure conotoxin peptide derivative comprising a

permutant of the peptide of claim 2.

Claim 12 (withdrawn): Use of a radiolabeled conotoxin peptide of claim 1 for

characterization of a new site on the aforementioned receptors or channels and use of these peptide

probes for screening and identification of novel small molecules that interact at the aforementioned

sites.

Claim 13 (withdrawn): The use of claim 12, wherein said receptor or channel is a

monoamine transporter.

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Claim 14 (withdrawn): The use of claim 13, wherein said peptide is selected from the group of peptides set forth in Table 5.

Claim 15 (new): The isolated peptide of claim 1 selected from the group consisting of:

ZTCCGYRMCVPC (SEQ ID NO:523);

ACCGYKLCSPC (SEQ ID NO:524);

STCCGFKMCIPCR (SEQ ID NO:525);

STCCGFKMCIPCS (SEQ ID NO:526);

STCCGFKMCIPC (SEQ ID NO:527);

STCCGYRMCVPC (SEQ ID NO:528);

NGVCCGYKLCLPC (SEQ ID NO:529);

LCCGFWMCIPCN (SEQ ID NO:530);

NGVCCGYKLCHOC (SEQ ID NO:531);

GVCCGYKLCHOC (SEQ ID NO:532);

SVCCGYKLCFPC (SEQ ID NO:534);

NGVCCGYRMCVPC (SEQ ID NO:535);

ZACCGFKMCVPC (SEQ ID NO:537);

NGVCCGFWMCIPCN (SEQ ID NO:539);

DVCCYVRMCPCR (SEQ ID NO:540); and

derivatives thereof.

Claim 16 (new): The isolated peptide of claim 15 selected from the group of:

SVCCGYKLCFPC (SEQ ID NO:534) and

derivatives thereof.

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Claim 17 (new): The isolated peptide of claim 16 that is SVCCGYKLCFPC (SEQ ID NO:534).

Claim 18 (new): The isolated conotoxin propertide of claim 6 selected from the group of conotoxin propertides of SEQ ID NO:62, SEQ ID NO:162, SEQ ID NO:229, SEQ ID NO:330, SEQ ID NO:333, SEQ ID NO:333, SEQ ID NO:336, SEQ ID NO:339, SEQ ID NO:342, SEQ ID NO:345, SEQ ID NO:352, SEQ ID NO:355, SEQ ID NO:361, SEQ ID NO:367 and SEQ ID NO:370.

Claim 19 (new): The isolated conotoxin propertide of claim 18 that is the conotoxin propertide of SEQ ID NO:352.